Characterization and Functional Implications of the NONEXPRESSOR OF PATHOGENESIS-RELATED GENES 1 (*NPR1*) in *Saccharum*

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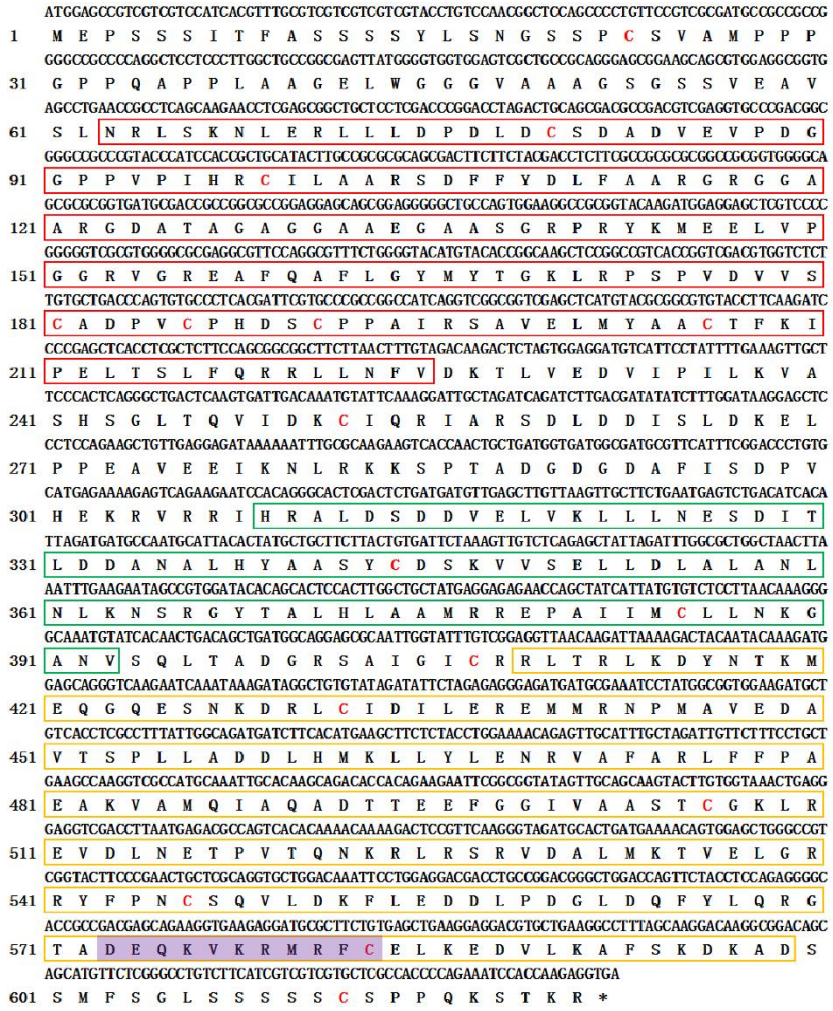

Figure. S1 Nucleic acid sequence and the coding amino acid sequence of *ShNPR1* gene. The red font was the cysteine site; the red box, green box and orange box sequences represented the BTB/POZ conservative domain, ankyrin repeats and NPR1-like C-terminal region, respectively. The purple sequence was the nuclear location signal, and * represented stop codon.